This listing of claims will replace all prior versions and listings of claims in the application.

LISTING OF CLAIMS

- 1-5. (Cancelled)
- 6. (Currently Amended) Surface carrying a linker system according to claim $\frac{121}{2}$.
- 7. (Currently Amended) Surface according to claim 6-A surface carrying a linker system comprising a compound for activating surfaces for bioconjugation having the following general formula (I):

$$X-[(Y_1)_i-Q-(Y_2)_i]_k-Z$$
 (I)

wherein:

X is a reactive group capable of covalently binding to a surface and is selected from the group consisting of a SiW₃ group with W being a hydrolyzable atom or group, an anthrathione group or a derivative thereof, an anthraquinone group or a derivative thereof, and a benzophenone group or a derivative thereof;

Z is a reactive group capable of covalently binding to a biomolecule, is capable of nucleophilic substitution reactions, nucleophilic addition reactions, Diels-Alder reactions or radical substitutions, and is selected from the group consisting of a diene group, a dienophilic group, an aldehyde group, a hydroxyl group, a carboxylic acid group, an active ester group, an amino group, a thiol group, an aziridine group, an isocyanate group, an isocyanate group, an azide group, and a reactive leaving group;

X is not Z;

 Y_1 and Y_2 are, independently from each other, CR_1R_2 ;

 R_1 and R_2 are, independently from each other, H, C_1 - C_4 alkyl, C_1 - C_4 alkoxy or C_1 - C_4 acyloxy;

i, j, and k are, independently from each other, an integer in the range from 1 to 10; the total number of C atoms in Y₁ and Y₂, the C atoms of R₁ and R₂ not included, is in the range of 2 to 100; Q is a hydrophilic atom or group selected from the group consisting of O, NH, C=O, O-C=O and CR₃R₄;

R₃ and R₄ are, independently from each other, selected from the group consisting of H, OH, C₁-C₄ alkoxy and C₁-C₄ acyloxy; and

R₃ and R₄ are not H at the same time;

wherein when Q = NH, Z is not NH_2 ;

wherein when k > 1, the Q's for each $[(Y_1)_{i}-Q-(Y_2)_{i}]_k$ are independently selected from each other; and

wherein said the linker system forms a patterned array.

- 8. (Previously Presented) Surface according to claim 6, wherein said surface is selected from the group consisting of a SiO₂ surface of a silicon wafer, glass, quartz, fused silica, gold and a polymer.
- 9. (Currently Amended) Surface according to any of claim 6, A surface carrying a linker system comprising a compound for activating surfaces for bioconjugation having the following general formula (I):

$$X-[(Y_1)_i-Q-(Y_2)_i]_k-Z$$
 (I)

___wherein:

X is a reactive group capable of covalently binding to a surface and is selected from the group consisting of a SiW₃ group with W being a hydrolyzable atom or group, an anthrathione group or a derivative thereof, an anthraquinone group or a derivative thereof, and a benzophenone group or a derivative thereof;

Z is a reactive group capable of covalently binding to a biomolecule, is capable of nucleophilic substitution reactions, nucleophilic addition reactions, Diels-Alder reactions or radical substitutions, and is selected from the group consisting of a diene group, a dienophilic group, an aldehyde group, a hydroxyl group, a carboxylic acid group, an active ester group, an amino group, a thiol group, an aziridine group, an isocyanate group, an isocyanate group, an azide group, and a reactive leaving group;

X is not Z;

 Y_1 and Y_2 are, independently from each other, CR_1R_2 ;

 R_1 and R_2 are, independently from each other, H, C_1 - C_4 alkyl, C_1 - C_4 alkoxy or C_1 - C_4 acyloxy;

i, j, and k are, independently from each other, an integer in the range from 1 to 10; the total number of C atoms in Y₁ and Y₂, the C atoms of R₁ and R₂ not included, is in the range of 2 to 100;

Q is a hydrophilic atom or group selected from the group consisting of O, NH, C=O, O-C=O and CR₃R₄;

 R_3 and R_4 are, independently from each other, selected from the group consisting of H, OH, C_1 - C_4 alkoxy and C_1 - C_4 acyloxy; and

R₃ and R₄ are not H at the same time;

wherein when Q = NH, Z is not NH_2 ;

wherein when k > 1, the Q's for each $[(Y_1)_{i-}Q_{-}(Y_2)_{j}]_k$ are independently selected from each other; and

wherein said the linker system is covalently bonded to a biomolecule.

- 10. (Original) Surface according to claim 9 wherein said biomolecule is a partner of a specifically interacting system of complementary binding partners.
- 11. (Original) Surface according to claim 10 wherein said specifically interacting system of complementary binding partners is based on nucleic acid/complementary nucleic acid, peptide nucleic acid/nucleic acid, enzyme/substrate, receptor/effector, lectin/sugar, antibody/antigen, avidin/biotin or streptavidin/biotin interaction.
- 12. (Original) Surface according to claim 11 wherein said nucleic acid is a DNA or RNA.
- 13. (Original) Surface according to claim 12 wherein said DNA or RNA is an oligonucleotide or an aptamer.

- 14. (Original) Surface according to claim 11 wherein said antibody is a polyclonal, monoclonal, chimeric or single-chain antibody or a functional fragment or derivative of such antibodies.
- 15. (Previously Presented) Process for the detection of a biomolecule which is a partner of a specifically interacting system of complementary binding partners, comprising the steps of:
- a) contacting a surface according to claim 10 with a sample suspected to contain the complementary binding partner,
- b) removing non-specifically bound sample components in a washing step, and
- c) detecting specifically bound sample components.
- 16. (Currently Amended) Process according to claim 15 wherein for said detecting, a colored, fluorescent, bioluminescent, chemoluminescent, phosporescent phosphorescent or radioactive label; an enzyme; an antibody or a functional fragment or derivative thereof, a protein A/gold based system; a biotin/avidin/streptavidin based system; or an enzyme electrode based system is used.
- 17. (Previously Presented) Process for the isolation of a biomolecule which is a partner of a specifically interacting system of complementary binding partners, comprising the steps of:
- a) contacting a surface according to claim 10 with a sample suspected to contain the biomolecule complementary binding partner,
- b) removing non-specifically bound sample components in a washing step, and, optionally,
- c) eluting specifically bound sample components.
- 18. (Previously Presented) A method of affinity chromatography comprising the steps of:

providing a surface according to claim 10 as an affinity matrix; and performing affinity chromatography with the affinity matrix.

- 19. (Previously Presented) A method of detecting a biomolecule comprising the steps of:providing a sensor chip or biochip comprising a surface according to claim 10; and detecting a biomolecule with the sensor chip or biochip.
- 20. (Previously Presented) Medical or diagnostic instrument comprising a surface according to claim 10.
- 21. (Currently Amended) A compound for activating surfaces for bioconjugation having the following general formula (I):

$$X-[(Y_1)_i-Q-(Y_2)_i]_k-Z$$
 (I)

wherein:

X is a reactive group capable of covalently binding to a surface and is selected from the group consisting of a SiW_3 group with W being a hydrolyzable atom or group, an anthrathione group or a derivative thereof, an anthraquinone group or a derivative thereof, and a benzophenone group or a derivative thereof;

Z is a reactive group capable of covalently binding to a biomolecule, is capable of nucleophilic substitution reactions, nucleophilic addition reactions, Diels-Alder reactions or radical substitutions, and is selected from the group consisting of a diene group, a dienophilic group, an aldehyde group, a hydroxyl group, a carboxylic acid group, an active ester group, an amino group, a thiol group, an aziridine group, an isocyanate group, an isotyanate group, an azide group, and a reactive leaving group;

X is not Z;

 Y_1 and Y_2 are, independently from each other, CR_1R_2 ;

 R_1 and R_2 are, independently from each other, H, C_1 - C_4 alkyl, C_1 - C_4 alkoxy or C_1 - C_4 acyloxy;

i, j, and k are, independently from each other, an integer in the range from 1 to 10; the total number of C atoms in Y₁ and Y₂, the C atoms of R₁ and R₂ not included, is in the range of 2 to 100;

Q is a hydrophilic atom or group selected from the group consisting of O, NH, C=O, O-C=O and CR₃R₄;

R₃ and R₄ are, independently from each other, selected from the group consisting of H, OH, C₁-C₄ alkoxy and C₁-C₄ acyloxy; and

R₃ and R₄ are not H at the same time;

wherein when Q = NH, Z is not NH_2 ;

wherein when k > 1, the Q's for each $[(Y_1)_{i-}Q_{-}(Y_2)_{j}]_k$ are independently selected from each other;

wherein when Z comprises a silane, X comprises a photocrosslinker; and wherein when Z comprises a photocrosslinker, X comprises a reactive group.

- 22. (Previously Presented) A process for the detection of a biomolecule, comprising the steps of:
- (a) providing a surface bound to a linker molecule in a patterned array, the linker molecule being covalently bound to a biomolecule,

the linker molecule having the following general formula (I):

$$X-[(Y_1)_i-Q-(Y_2)_i]_k-Z$$
 (I)

wherein:

X is a reactive group capable of covalently binding to a surface and is selected from the group consisting of a SiW₃ group with W being a hydrolyzable atom or group, an anthrathione group or a derivative thereof, an anthraquinone group or a derivative thereof, and a benzophenone group or a derivative thereof;

Z is a reactive group capable of covalently binding to a biomolecule, is capable of nucleophilic substitution reactions, nucleophilic addition reactions, Diels-Alder reactions or radical substitutions, and is selected from the group consisting of a diene group, a dienophilic group, an aldehyde group, a hydroxyl group, a carboxylic acid group, an active ester group, an amino group, a thiol group, an aziridine group, an isocyanate group, an isothiocyanate group, an azide group, and a reactive leaving group;

X is not Z;

 Y_1 and Y_2 are, independently from each other, CR_1R_2 ;

 R_1 and R_2 are, independently from each other, H, C_1 - C_4 alkyl, C_1 - C_4 alkoxy or C_1 - C_4 acyloxy;

i, j, and k are, independently from each other, an integer in the range from 1 to 10;

the total number of C atoms in Y_1 and Y_2 , the C atoms of R_1 and R_2 not included, is in the range of 2 to 100;

Q is a hydrophilic atom or group selected from the group consisting of O, NH, C=O, O-C=O and CR₃R₄;

R₃ and R₄ are, independently from each other, selected from the group consisting of H, OH, C₁-C₄ alkoxy and C₁-C₄ acyloxy; and

R₃ and R₄ are not H at the same time;

wherein when Q = NH, Z is not NH_2 ;

wherein when k > 1, the Q's for each $[(Y_1)_i-Q-(Y_2)_j]_k$ are independently selected from each other; and

wherein the biomolecule is a partner of one or more specifically interacting complementary binding partners based on nucleic acid/complementary nucleic acid, peptide nucleic acid/nucleic acid, enzyme/substrate, receptor/effector, lectin/sugar, antibody/antigen, avidin/biotin or streptavidin/biotin interaction;

- (b) contacting the surface with a sample to be tested;
- (c) removing non-specifically bound sample components in a washing step; and
- (d) detecting specifically bound sample components.
- 23. (Previously Presented) The method of claim 22, wherein said surface comprises a silicon oxide or gold.